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			420					425						430	
Leu	Val	Asp	Gln	Thr	Glu	Gly	Cys	Gln	Pro	Val	Tyr	Val			
		435					440					445			

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<400>	16															
Met	Phe	Cys	Ser	Phe	Glu	Thr	Arg	Arg	His	Leu	Cys	Met	Val	Met	Glu	
1				5					10					15		
Tyr	Val	Glu	Gly	Gly	Asp	Cys	Ala	Thr	Leu	Leu	Lys	Asn	Ile	Gly	Ala	
			20					25					30			
Leu	Pro	Val	Glu	Met	Ala	Arg	Met	Tyr	Phe	Ala	Glu	Thr	Val	Leu	Ala	
		35					40					45				
Leu	Glu	Tyr	Leu	His	Asn	Tyr	Gly	Ile	Val	His	Arg	Asp	Leu	Lys	Pro	
	50					55					60					
Asp	Asn	Leu	Leu	Ile	Thr	Ser	Met	Gly	His	Ile	Lys	Leu	Thr	Asp	Phe	
65					70					75					80	
Gly	Leu	Ser	Lys	Met	Gly	Leu	Met	Ser	Leu	Thr	Thr	Asn	Leu	Tyr	Glu	
				85					90					95		
Gly	His	Ile	Glu	Lys	Asp	Ala	Arg	Glu	Phe	Leu	Asp	Lys	Gln	Val	Cys	
			100					105					110			
Gly	Thr	Pro	Glu	Tyr	Ile	Ala	Pro	Glu	Val	Ile	Leu	Arg	Gln	Gly	Tyr	
		115				120						125				
Gly	Lys	Pro	Val	Asp	Trp	Trp	Ala	Met	Gly	Ile	Ile	Leu	Tyr	Glu	Phe	
	130					135					140					
Leu	Val	Gly	Cys	Val	Pro	Phe	Phe	Gly	Asp	Thr	Pro	Glu	Glu	Leu	Phe	
145					150				155						160	
Gly	Gln	Val	Ile	Ser	Asp	Asp	Ile	Leu	Trp	Pro	Glu	Gly	Asp	Glu	Ala	
				165					170					175		
Leu	Pro	Thr	Glu	Ala	Gln	Leu	Leu	Ile	Ser	Ser	Leu	Leu	Gln	Thr	Asn	
			180					185					190			
Pro	Leu	Val	Arg	Leu	Gly	Ala	Gly	Gly	Ala	Phe	Glu	Val	Lys	Gln	His	
		195				200						205				
Ser	Phe	Phe	Arg	Asp	Leu	Asp	Trp	Thr	Gly	Leu	Leu	Arg	Gln	Lys	Ala	
	210					215					220					
Glu	Phe	Ile	Pro	His	Leu	Glu	Ser	Glu	Asp	Asp	Thr	Ser	Tyr	Phe	Asp	
225					230				235						240	
Thr	Arg	Ser	Asp	Arg	Tyr	His	His	Val	Asn	Ser	Tyr	Asp	Glu	Asp	Asp	
				245					250					255		
Thr	Thr	Glu	Glu	Glu	Pro	Val	Glu	Ile	Arg	Gln	Phe	Ser	Ser	Cys	Ser	
			260					265					270			

Pro	Arg	Phe	Ser	Lys	Val	Tyr	Ser	Ser	Met	Glu	Gln	Leu	Ser	Gln	His
		275					280					285			
Glu	Pro	Lys	Thr	Pro	Val	Ala	Ala	Ala	Gly	Ser	Ser	Lys	Arg	Glu	Pro
	290					295					300				
Ser	Thr	Lys	Gly	Pro	Glu	Glu	Lys	Val	Ala	Gly	Lys	Arg	Glu	Gly	Leu
305					310					315					320
Gly	Gly	Leu	Thr	Leu	Arg	Glu	Lys	Thr	Trp	Arg	Gly	Gly	Ser	Pro	Glu
				325					330					335	
Ile	Lys	Arg	Phe	Ser	Ala	Ser	Glu	Ala	Ser	Phe	Leu	Glu	Gly	Glu	Ala
			340					345					350		
Ser	Pro	Pro	Leu	Gly	Ala	Arg	Arg	Arg	Phe	Ser	Ala	Leu	Leu	Glu	Pro
		355					360					365			
Ser	Arg	Phe	Ser	Ala	Pro	Gln	Glu	Asp	Glu	Asp	Glu	Ala	Arg	Leu	Arg
	370					375					380				
Arg	Pro	Pro	Arg	Pro	Ser	Ser	Asp	Pro	Ala	Gly	Ser	Leu	Asp	Ala	Arg
385					390					395					400
Ala	Pro	Lys	Glu	Glu	Thr	Gln	Gly	Glu	Gly	Thr	Ser	Ser	Ala	Gly	Asp
				405					410					415	
Ser	Glu	Ala	Thr	Asp	Arg	Pro	Arg	Pro	Gly	Asp	Leu	Cys	Pro	Pro	Ser
			420					425					430		
Lys	Asp	Gly	Asp	Ala	Ser	Gly	Pro	Arg	Ala	Thr	Asn	Asp	Leu	Val	Leu
		435					440					445			
Arg	Arg	Ala	Arg	His	Gln	Gln	Met	Ser	Gly	Asp	Val	Ala	Val	Glu	Lys
		450				455					460				
Arg	Pro	Ser	Arg	Thr	Gly	Gly	Lys	Val	Ile	Lys	Ser	Ala	Ser	Ala	Thr
465					470					475					480
Ala	Leu	Ser	Val	Met	Ile	Pro	Ala	Val	Asp	Pro	His	Gly	Ser	Ser	Pro
				485					490					495	
Leu	Ala	Ser	Pro	Met	Ser	Pro	Arg	Ser	Leu	Ser	Ser	Asn	Pro	Ser	Ser
			500					505					510		
Arg	Asp	Ser	Ser	Pro	Ser	Arg	Asp	Tyr	Ser	Pro	Ala	Val	Ser	Gly	Leu
		515					520					525			
Arg	Ser	Pro	Ile	Thr	Ile	Gln	Arg	Ser	Gly	Lys	Lys	Tyr	Gly	Phe	Thr
		530				535					540				
Leu	Arg	Ala	Ile	Arg	Val	Tyr	Met	Gly	Asp	Thr	Asp	Val	Tyr	Ser	Val
545					550					555					560
His	His	Ile	Val	Trp	His	Val	Glu	Glu	Gly	Gly	Pro	Ala	Gln	Glu	Ala
				565					570					575	

Gly	Leu	Cys	Ala 580	Gly	Asp	Leu	Ile	Thr 585	His	Val	Asn	Gly	Glu 590	Pro	Val
His	Gly	Met 595	Val	His	Pro	Glu	Val 600	Val	Glu	Leu	Ile	Leu 605	Lys	Ser	Gly
Asn	Lys 610	Val	Ala	Val	Thr	Thr 615	Thr	Pro	Phe	Glu	Asn 620	Thr	Ser	Ile	Arg
Ile 625	Gly	Pro	Ala	Arg	Arg 630	Ser	Ser	Tyr	Lys	Ala 635	Lys	Met	Ala	Arg	Arg 640
Asn	Lys	Arg	Pro	Ser 645	Ala	Lys	Glu	Gly	Gln 650	Glu	Ser	Lys	Lys	Arg 655	Ser
Ser	Leu	Phe 660	Arg	Lys	Ile	Thr	Lys	Gln 665	Ser	Asn	Leu	Leu	His 670	Thr	Ser
Arg	Ser	Leu 675	Ser	Ser	Leu	Asn	Arg 680	Ser	Leu	Ser	Ser	Ser 685	Asp	Ser	Leu
Pro	Gly 690	Ser	Pro	Thr	His	Gly 695	Leu	Pro	Ala	Arg	Ser 700	Pro	Thr	His	Ser
Tyr 705	Arg	Ser	Thr	Pro	Asp 710	Ser	Ala	Tyr	Leu	Gly 715	Ala	Ser	Ser	Gln	Ser 720
Ser	Ser	Pro	Ala	Ser 725	Ser	Thr	Pro	Asn	Ser 730	Pro	Ala	Ser	Ser	Ala	Ser 735
His	His	Ile 740	Arg	Pro	Ser	Thr	Leu	His 745	Gly	Leu	Ser	Pro	Lys 750	Leu	His
Arg	Gln	Tyr 755	Arg	Ser	Ala	Arg	Cys 760	Lys	Ser	Ala	Gly	Asn 765	Ile	Pro	Leu
Ser	Pro 770	Leu	Ala	His	Thr	Pro 775	Ser	Pro	Thr	Gln	Ala 780	Ser	Pro	Pro	Pro
Leu 785	Pro	Gly	His	Thr 790	Val	Gly	Ser	Ser	His	Thr 795	Thr	Gln	Ser	Phe	Pro 800
Ala	Lys	Leu	His	Ser 805	Ser	Pro	Pro	Val	Val 810	Arg	Pro	Arg	Pro	Lys 815	Ser
Ala	Glu	Pro	Pro 820	Arg	Ser	Pro	Leu	Leu 825	Lys	Arg	Val	Gln	Ser 830	Ala	Glu
Lys	Leu	Gly 835	Ala	Ser	Leu	Ser	Ala 840	Asp	Lys	Lys	Gly	Ala 845	Leu	Arg	Lys
His 850	Ser	Leu	Glu	Val	Gly	His 855	Pro	Asp	Phe	Arg	Lys 860	Asp	Phe	His	Gly
Glu 865	Leu	Ala	Leu	His	Ser 870	Leu	Ala	Glu	Ser	Asp 875	Gly	Glu	Thr	Pro	Pro 880
Val	Glu	Gly	Leu	Gly	Ala	Pro	Arg	Gln	Val	Ala	Val	Arg	Arg	Leu	Gly

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<210> 17
<211> 356
<212> PRT
<213> Homo sapiens

<400> 17
Met Lys Ser Arg Val Lys Gly Ser Pro Val Tyr Thr Ala Pro Glu Val
 1             5             10             15
Val Arg Gly Ala Asp Phe Ser Ile Ser Ser Asp Leu Trp Ser Leu Gly
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<400> 19  
Met Ile His Arg Asp Leu Lys Pro Val Asn Ile Phe Leu Asp Ser Asp

1	5	10	15
Asp His Val Lys Ile Gly Asp Phe Gly Leu Ala Thr Asp His Leu Ala	20	25	30
Phe Ser Ala Asp Ser Lys Gln Asp Asp Gln Thr Gly Asp Leu Ile Lys	35	40	45
Ser Asp Pro Ser Gly His Leu Thr Gly Met Val Gly Thr Ala Leu Tyr	50	55	60
Val Ser Pro Glu Val Gln Gly Ser Thr Lys Ser Ala Tyr Asn Gln Lys	65	70	75
Val Asp Leu Phe Ser Leu Gly Ile Ile Phe Phe Glu Met Ser Tyr His	85	90	95
Pro Met Val Thr Ala Ser Glu Arg Ile Phe Val Leu Asn Gln Leu Arg	100	105	110
Asp Pro Thr Ser Pro Lys Phe Pro Glu Asp Phe Asp Asp Gly Glu His	115	120	125
Ala Lys Gln Lys Ser Val Ile Ser Trp Leu Leu Asn His Asp Pro Ala	130	135	140
Lys Arg Pro Thr Ala Thr Glu Leu Leu Lys Ser Glu Leu Leu Pro Pro	145	150	155
Pro Gln Met Glu Glu Ser Glu Leu His Glu Val Leu His His Thr Leu	165	170	175
Thr Asn Val Asp Gly Lys Ala Tyr Arg Thr Met Met Ala Gln Ile Phe	180	185	190
Ser Gln Arg Ile Ser Pro Ala Ile Asp Tyr Thr Tyr Asp Ser Asp Ile	195	200	205
Leu Lys Gly Asn Phe Ser Ile Arg Thr Ala Lys Met Gln Gln His Val	210	215	220
Cys Glu Thr Ile Ile Arg Ile Phe Lys Arg His Gly Ala Val Gln Leu	225	230	235
Cys Thr Pro Leu Leu Leu Pro Arg Asn Arg Gln Ile Tyr Glu His Asn	245	250	255
Glu Ala Ala Leu Phe Met Asp His Ser Gly Met Leu Val Met Leu Pro	260	265	270
Phe Asp Leu Arg Ile Pro Phe Ala Arg Tyr Val Ala Arg Asn Asn Ile	275	280	285
Leu Asn Leu Lys Arg Tyr Cys Ile Glu Arg Val Phe Arg Pro Arg Lys	290	295	300
Leu Asp Arg Phe His Pro Lys Glu Leu Leu Glu Cys Ala Phe Asp Ile	305	310	315
			320

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Val	Thr	Ser	Thr	Thr	Asn	Ser	Phe	Leu	Pro	Thr	Ala	Glu	Ile	Ile	Tyr	
				325					330					335		
Thr	Ile	Tyr	Glu	Ile	Ile	Gln	Glu	Phe	Pro	Ala	Leu	Gln	Glu	Arg	Asn	
				340					345					350		
Tyr	Ser	Ile	Tyr	Leu	Asn	His	Thr	Met	Leu	Leu	Lys	Ala	Ile	Leu	Leu	
				355					360					365		
His	Cys	Gly	Ile	Pro	Glu	Asp	Lys	Leu	Ser	Gln	Val	Tyr	Ile	Ile	Leu	
				370					375					380		
Tyr	Asp	Ala	Val	Thr	Glu	Lys	Leu	Thr	Arg	Arg	Glu	Val	Glu	Ala	Lys	
				385					390					395		
Phe	Cys	Asn	Leu	Ser	Leu	Ser	Ser	Asn	Ser	Leu	Cys	Arg	Leu	Tyr	Lys	
				405					410					415		
Phe	Ile	Glu	Gln	Lys	Gly	Asp	Leu	Gln	Asp	Leu	Met	Pro	Thr	Ile	Asn	
				420					425					430		
Ser	Leu	Ile	Lys	Gln	Lys	Thr	Gly	Ile	Ala	Gln	Leu	Val	Lys	Tyr	Gly	
				435					440					445		
Leu	Lys	Asp	Leu	Glu	Glu	Val	Val	Gly	Leu	Leu	Lys	Lys	Leu	Gly	Ile	
				450					455					460		
Lys	Leu	Gln	Val	Leu	Ile	Asn	Leu	Gly	Leu	Val	Tyr	Lys	Val	Gln	Gln	
				465					470					475		
His	Asn	Gly	Ile	Ile	Phe	Gln	Phe	Val	Ala	Phe	Ile	Lys	Arg	Arg	Gln	
				485					490					495		
Arg	Ala	Val	Pro	Glu	Ile	Leu	Ala	Ala	Gly	Gly	Arg	Tyr	Asp	Leu	Leu	
				500					505					510		
Ile	Pro	Gln	Phe	Arg	Gly	Pro	Gln	Ala	Leu	Gly	Pro	Val	Pro	Thr	Ala	
				515					520					525		
Ile	Gly	Val	Ser	Ile	Ala	Ile	Asp	Lys	Ile	Ser	Ala	Ala	Val	Leu	Asn	
				530					535					540		
Met	Glu	Glu	Ser	Val	Thr	Ile	Ser	Ser	Cys	Asp	Leu	Leu	Val	Val	Ser	
				545					550					555		
Val	Gly	Gln	Met	Ser	Met	Ser	Arg	Ala	Ile	Asn	Leu	Thr	Gln	Lys	Leu	
				565					570					575		
Trp	Thr	Ala	Gly	Ile	Thr	Ala	Glu	Ile	Met	Tyr	Asp	Trp	Ser	Gln	Ser	
				580					585					590		
Gln	Glu	Glu	Leu	Gln	Glu	Tyr	Cys	Arg	His	His	Glu	Ile	Thr	Tyr	Val	
				595					600					605		
Ala	Leu	Val	Ser	Asp	Lys	Glu	Gly	Ser	His	Val	Lys	Val	Lys	Ser	Phe	
				610					615					620		

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<210> 20
<211> 530
<212> PRT
<213> Homo sapiens

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Arg Arg Leu Ser Glu Arg Glu Thr Arg His Phe Phe Arg Gln Ile Val
          20             25             30
Ser Ala Val His Tyr Cys His Lys Asn Gly Val Val His Arg Asp Leu
          35             40             45
Lys Leu Glu Asn Ile Leu Leu Asp Asp Asn Cys Asn Ile Lys Ile Ala
 50             55             60
Asp Phe Gly Leu Ser Asn Leu Tyr Gln Lys Asp Lys Phe Leu Gln Thr
 65             70             75             80

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Phe	Cys	Gly	Ser	Pro 85	Leu	Tyr	Ala	Ser	Pro 90	Glu	Ile	Val	Asn	Gly 95	Arg
Pro	Tyr	Arg	Gly 100	Pro	Glu	Val	Asp	Ser 105	Trp	Ala	Leu	Gly	Val 110	Leu	Leu
Tyr	Thr	Leu 115	Val	Tyr	Gly	Thr	Met 120	Pro	Phe	Asp	Gly	Phe 125	Asp	His	Lys
Asn	Leu 130	Ile	Arg	Gln	Ile	Ser 135	Ser	Gly	Glu	Tyr	Arg 140	Glu	Pro	Thr	Gln
Pro 145	Ser	Asp	Ala	Arg	Gly 150	Leu	Ile	Arg	Trp	Met 155	Leu	Met	Val	Asn	Pro 160
Asp	Arg	Arg	Ala	Thr 165	Ile	Glu	Asp	Ile	Ala 170	Asn	His	Trp	Trp	Val 175	Asn
Trp	Gly	Tyr	Lys 180	Ser	Ser	Val	Cys	Asp 185	Cys	Asp	Ala	Leu	His 190	Asp	Ser
Glu	Ser 195	Pro	Leu	Leu	Ala	Arg	Ile 200	Ile	Asp	Trp	His	His 205	Arg	Ser	Thr
Gly 210	Leu	Gln	Ala	Asp	Thr	Glu 215	Ala	Lys	Met	Lys	Gly 220	Leu	Ala	Lys	Pro
Thr 225	Thr	Ser	Glu	Val 230	Met	Leu	Glu	Arg	Gln 235	Arg	Ser	Leu	Lys	Lys	Ser 240
Lys	Lys	Glu	Asn 245	Asp	Phe	Ala	Gln	Ser	Gly 250	Gln	Asp	Ala	Val	Pro 255	Glu
Ser	Pro	Ser	Lys 260	Leu	Ser	Ser	Lys	Arg 265	Pro	Lys	Gly	Ile	Leu	Lys	Lys
Arg	Ser 275	Asn	Ser	Glu	His	Arg	Ser 280	His	Ser	Thr	Gly	Phe 285	Ile	Glu	Gly
Val 290	Val	Gly	Pro	Ala	Leu	Pro 295	Ser	Thr	Phe	Lys	Met 300	Glu	Gln	Asp	Leu
Cys 305	Arg	Thr	Gly	Val 310	Leu	Leu	Pro	Ser	Ser	Pro 315	Glu	Ala	Glu	Val	Pro 320
Gly	Lys	Leu	Ser 325	Pro	Lys	Gln	Ser	Ala 330	Thr	Met	Pro	Lys	Lys	Gly 335	Ile
Leu	Lys	Lys	Thr 340	Gln	Gln	Arg	Glu	Ser 345	Gly	Tyr	Tyr	Ser	Ser 350	Pro	Glu
Arg	Ser 355	Glu	Ser	Ser	Glu	Leu	Leu 360	Asp	Ser	Asn	Asp	Val 365	Met	Gly	Ser
Ser 370	Ile	Pro	Ser	Pro	Ser	Pro 375	Pro	Asp	Pro	Ala	Arg 380	Val	Thr	Ser	His
Ser	Leu	Ser	Cys	Arg	Arg	Lys	Gly	Ile	Leu	Lys	His	Ser	Ser	Lys	Tyr

385                      390                      395                      400  
 Ser Ala Gly Thr Met Asp Pro Ala Leu Val Ser Arg Glu Met Pro Thr  
                                  405                      410                      415  
 Leu Glu Ser Leu Ser Glu Pro Gly Val Pro Ala Glu Gly Leu Ser Arg  
                                  420                      425                      430  
 Ser Tyr Ser Arg Pro Ser Ser Val Ile Ser Asp Asp Ser Val Leu Ser  
                                  435                      440                      445  
 Ser Asp Ser Phe Asp Leu Leu Asp Leu Gln Glu Asn Arg Pro Ala Arg  
                                  450                      455                      460  
 Gln Arg Ile Arg Ser Cys Val Ser Ala Glu Asn Phe Leu Gln Ile Gln  
                                  465                      470                      475                      480  
 Asp Phe Glu Gly Leu Gln Asn Arg Pro Arg Pro Gln Tyr Leu Lys Arg  
                                  485                      490                      495  
 Tyr Arg Asn Arg Leu Ala Asp Ser Ser Phe Ser Leu Leu Thr Asp Met  
                                  500                      505                      510  
 Asp Asp Val Thr Gln Val Tyr Lys Gln Ala Leu Glu Ile Cys Ser Lys  
                                  515                      520                      525  
 Leu Asn  
                                  530  
  
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 <211> 728  
 <212> PRT  
 <213> Homo sapiens  
  
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 Phe Arg Gln Glu Ala Ser Met Leu His Ala Leu Gln His Pro Cys Ile  
                                   20                                  25                                  30  
 Val Ala Leu Ile Gly Ile Ser Ile His Pro Leu Cys Phe Ala Leu Glu  
                                   35                                  40                                  45  
 Leu Ala Pro Leu Ser Ser Leu Asn Thr Val Leu Ser Glu Asn Ala Arg  
                                   50                                  55                                  60  
 Asp Ser Ser Phe Ile Pro Leu Gly His Met Leu Thr Gln Lys Ile Ala  
                                   65                                  70                                  75                                  80  
 Tyr Gln Ile Ala Ser Gly Leu Ala Tyr Leu His Lys Lys Asn Ile Ile  
                                   85                                  90                                  95  
 Phe Cys Asp Leu Lys Ser Asp Asn Ile Leu Val Trp Ser Leu Asp Val  
                                   100                                  105                                  110  
 Lys Glu His Ile Asn Ile Lys Leu Ser Asp Tyr Gly Ile Ser Arg Gln

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115	120	125
Ser Phe His Glu Gly Ala Leu Gly Val Glu Gly Thr Pro Gly Tyr Gln		
130	135	140
Ala Pro Glu Ile Arg Pro Arg Ile Val Tyr Asp Glu Lys Val Asp Met		
145	150	155
Phe Ser Tyr Gly Met Val Leu Tyr Glu Leu Leu Ser Gly Gln Arg Pro		
165	170	175
Ala Leu Gly His His Gln Leu Gln Ile Ala Lys Lys Leu Ser Lys Gly		
180	185	190
Ile Arg Pro Val Leu Gly Gln Pro Glu Glu Val Gln Phe Arg Arg Leu		
195	200	205
Gln Ala Leu Met Met Glu Cys Trp Asp Thr Lys Pro Glu Lys Arg Pro		
210	215	220
Leu Ala Leu Ser Val Val Ser Gln Met Lys Asp Pro Thr Phe Ala Thr		
225	230	235
Phe Met Tyr Glu Leu Cys Cys Gly Lys Gln Thr Ala Phe Phe Ser Ser		
245	250	255
Gln Gly Gln Glu Tyr Thr Val Val Phe Trp Asp Gly Lys Glu Glu Ser		
260	265	270
Arg Asn Tyr Thr Val Val Asn Thr Glu Lys Gly Leu Met Glu Val Gln		
275	280	285
Arg Met Cys Cys Pro Gly Met Lys Val Ser Cys Gln Leu Gln Val Gln		
290	295	300
Arg Ser Leu Trp Thr Ala Thr Glu Asp Gln Lys Ile Tyr Ile Tyr Thr		
305	310	315
Leu Lys Gly Met Cys Pro Leu Asn Thr Pro Gln Gln Ala Leu Asp Thr		
325	330	335
Pro Ala Val Val Thr Cys Phe Leu Ala Val Pro Val Ile Lys Lys Asn		
340	345	350
Ser Tyr Leu Val Leu Ala Gly Leu Ala Asp Gly Leu Val Ala Val Phe		
355	360	365
Pro Val Val Arg Gly Thr Pro Lys Asp Ser Cys Ser Tyr Leu Cys Ser		
370	375	380
His Thr Ala Asn Arg Ser Lys Phe Ser Ile Ala Asp Glu Asp Ala Arg		
385	390	395
Gln Asn Pro Tyr Pro Val Lys Ala Met Glu Val Val Asn Ser Gly Ser		
405	410	415
Glu Val Trp Tyr Ser Asn Gly Pro Gly Leu Leu Val Ile Asp Cys Ala		
420	425	430

093639-041001



[illegible]

<210> 22  
 <211> 253  
 <212> PRT  
 <213> Homo sapiens

<400> 22  
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 Asp Gly Val Val His His His Gly Leu Phe Gln Asp Arg Thr Cys Glu  
                   20                  25                  30  
 Ile Val Glu Asp Thr Glu Ser Ser Arg Met Val Lys Lys Met Lys Lys  
           35                  40                  45  
 Arg Ile Cys Leu Val Leu Asp Cys Leu Cys Ala His Asp Phe Ser Asp  
       50                  55                  60  
 Lys Thr Ala Asp Leu Ile Asn Leu Gln His Tyr Val Ile Lys Glu Lys  
       65                  70                  75                  80  
 Arg Leu Ser Glu Arg Glu Thr Val Val Ile Phe Tyr Asp Val Val Arg  
                   85                  90                  95  
 Val Val Glu Ala Leu His Gln Lys Asn Ile Val His Arg Asp Leu Lys  
           100                  105                  110  
 Leu Gly Asn Met Val Leu Asn Lys Arg Thr His Arg Ile Thr Ile Thr  
           115                  120                  125  
 Asn Phe Cys Leu Gly Lys His Leu Val Ser Glu Gly Asp Leu Leu Lys  
       130                  135                  140  
 Asp Gln Arg Gly Ser Pro Ala Tyr Ile Ser Pro Asp Gly Ser Ala Ala  
       145                  150                  155                  160  
 Gly Arg Thr Val Ala Ser Pro Val Thr Cys Gly Pro Trp Ala Trp Cys  
           165                  170                  175  
 Ser Ser Pro Cys Cys Met Ala Ser Ser Pro Ser Thr Thr Ala Ser Arg  
           180                  185                  190  
 Arg Ser Ser Ser Ala Arg Ser Arg Leu Pro Ser Ile Pro Phe Leu Arg  
       195                  200                  205  
 Met Asp Gly Phe Leu Arg Thr Pro Cys Val Ser Ser Gly Ser Cys Trp  
       210                  215                  220  
 Ser Leu Thr Pro Ser Ser Ala Trp Pro Pro Pro Thr Ser Trp Arg Pro  
       225                  230                  235                  240  
 Ser Val Pro Ser Leu His His Gly Ser Pro Cys His Leu  
           245                  250

<210> 23  
 <211> 131

05030302 0413001

1. The first part of the document discusses the importance of maintaining accurate records of all transactions and activities. It emphasizes that proper record-keeping is essential for transparency and accountability, particularly in financial matters.

2. The second part outlines the specific procedures for handling sensitive information. It stresses the need for strict confidentiality protocols to protect data from unauthorized access or disclosure.

3. The third part addresses the role of communication in ensuring the success of the project. It highlights the importance of clear, concise, and timely communication between all stakeholders involved.

4. The fourth part discusses the importance of regular monitoring and evaluation. It notes that ongoing assessment is necessary to identify potential issues early and make adjustments as needed.

5. The fifth part concludes by reiterating the commitment to high standards of performance and integrity. It expresses confidence in the team's ability to achieve the desired outcomes.

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<400> 24
Met Val Thr Leu Gly Ala Pro Pro Tyr Pro Glu Val Pro Pro Thr Ser
  1                               10                          15

Ile Leu Glu His Leu Gln Arg Arg Lys Ile Met Lys Arg Pro Ser Ser
      20                      25                      30

Cys Thr His Thr Met Tyr Ser Ile Met Lys Ser Cys Trp Arg Trp Arg
      35                      40                      45

Glu Ala Asp Arg Pro Ser Pro Arg Glu Leu Arg Leu Arg Leu Glu Ala
  50                      55                      60

Ala Ile Lys Thr Ala Asp Asp Glu Ala Val Leu Gln Val Pro Glu Leu
  65                      70                      75                      80

Val Val Pro Glu Leu Tyr Ala Ala Val Ala Gly Ile Arg Val Glu Ser
      85                      90                      95

Leu Phe Tyr Asn Tyr Ser Met Leu
      100

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```

<400> 25
Met Thr Gln Glu Val Val Thr Gln Tyr Tyr Arg Ala Pro Glu Ile Leu
  1                               10                      15

Met Gly Ser Arg His Tyr Ser Asn Ala Ile Asp Ile Trp Ser Val Gly
  20                      25                      30

Cys Ile Phe Ala Glu Leu Leu Gly Arg Arg Ile Leu Phe Gln Ala Gln
  35                      40                      45

Ser Pro Ile Gln Gln Leu Asp Leu Ile Thr Asp Leu Leu Gly Thr Pro
  50                      55                      60

Ser Leu Glu Ala Met Arg Thr Ala Cys Glu Gly Ala Lys Ala His Ile
  65                      70                      75                      80

Leu Arg Gly Pro His Lys Gln Val Arg Gly Gly Gly Asn Leu Phe Leu
  85                      90                      95

Val Thr Ile Cys Leu Gly Lys Ile Ser Leu Pro Ile Glu
  100                      105

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<400> 26																
Gly	Thr	Arg	Asp	Lys	Lys	Leu	Trp	Arg	Thr	Phe	Phe	Ile	Ser	Pro	Ala	
1				5					10					15		
Asn	Asn	Ile	Cys	Phe	Tyr	Gly	Glu	Cys	Ser	Tyr	Tyr	Cys	Ser	Thr	Glu	
			20					25					30			
His	Ala	Leu	Cys	Gly	Lys	Pro	Asp	Gln	Ile	Glu	Gly	Ser	Leu	Ala	Ala	
		35					40					45				
Phe	Leu	Pro	Asp	Leu	Ser	Leu	Ala	Lys	Arg	Lys	Thr	Trp	Arg	Asn	Pro	
	50					55					60					
Trp	Arg	Arg	Ser	Tyr	His	Lys	Arg	Lys	Lys	Ala	Glu	Trp	Glu	Val	Asp	
65					70					75					80	
Pro	Asp	Tyr	Cys	Glu	Glu	Val	Lys	Gln	Thr	Pro	Pro	Tyr	Asp	Ser	Ser	
				85					90					95		
His	Arg	Ile	Leu	Asp	Val	Met	Asp	Met	Thr	Ile	Phe	Asp	Phe	Leu	Met	
			100					105					110			
Gly	Asn	Met	Asp	Arg	His	His	Tyr	Glu	Thr	Phe	Glu	Lys	Phe	Gly	Asn	

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<210> 27
<211> 70
<212> PRT
<213> Homo sapiens

<400> 27
Met Tyr Asn Ser Val Lys Gly Ser Cys Ser Glu Pro Val Ser Phe Thr
  1              5              10              15
Thr His Ser Cys Ala Pro Glu Cys Pro Phe Pro Pro Lys Leu Ala His
          20              25              30
Arg Ser Lys Ser Ser Leu Thr Leu Gln Trp Lys Ala Pro Ile Asp Asn
          35              40              45
Gly Ser Lys Ile Thr Asn Tyr Leu Leu Glu Trp Asp Glu Val Ser Leu
  50              55              60
Phe Ser Tyr Ser Pro Ile
  65              70

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<210> 28
<211> 15
<212> PRT
<213> Homo sapiens

<400> 28
Lys Ile Lys Thr Leu Gly Ile Gly Ala Phe Gly Glu Val Cys Leu
 1           5           10          15
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<400> 32  
Gly Thr Ser Ile Leu Gly Pro Arg Leu Gly Asn Ser Pro Val Pro Ser  
1 5 10 15  
Ile Val Gln Cys Leu Ala Arg Lys Asp Gly Thr Asp Asp Phe Tyr Gln  
20 25 30

His